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| **Supplementary Table 2. Peptides used to Identify LCN2 Protein by MS** | | | | |
| **No.** | **Sequence** | **PEP** | **Charge** | **Molecular Weight (Da)** |
| 1 | VPLQQNFQDNQFQGK | 9.224E-11 | 2 | 1790.88 |
| 2 | MYATIYELKEDK | 2.411E-08 | 3 | 1503.74 |
| 3 | SYPGLTSYLVR | 0.0004464 | 2 | 1255.66 |
| 4 | VVSTNYNQHAMVFFKK | 0.00118 | 4 | 1928.96 |
| 5 | ELTSELKENFIR | 0.000009 | 3 | 1478.78 |
| PEP: value represents the reliability of mass spectrometric identification of peptide segments, usually the standard is 0.1 or 0.5, and the lower the value, the higher the reliability of peptide segment identification. | | | | |